

AMENDED LISTING OF CLAIMS

This listing of claims will replace all prior versions, and listings, of claims in this application:

1. (Currently amended) A method for determining resonant frequencies of electromagnetic radiation emission for influencing a medium surrounding a target genomic material nucleic acid chain, the genomic material said nucleic acid chain surrounded by a medium being sensitive to the electromagnetic response characteristics of the surrounding medium, comprising:
 - providing a frequency-emitting device capable of producing a frequency-influenced electric field, or magnetic field, or electromagnetic field, or electrical current emission;
 - determining a velocity for the propagation of the electromagnetic radiation emission through the medium surrounding the genomic material target nucleic acid chain;
 - determining a wavelength length parameter of the genomic material target nucleic acid chain when said target nucleic acid chain material consists of double-stranded or single-stranded molecules consisting of deoxyribonucleic acid or ribonucleic acid; the genomic material said target nucleic acid chain comprising a plurality of base pairs nucleotide bases spaced apart by an average spacing, the average spacing comprising a known value, by determining obtaining the number of base pairs nucleotide bases in the genomic material in a single strand of the target nucleic acid chain, in the case of double-stranded molecules not including the number of nucleotide bases in the complementary strand; and multiplying the said number of base pairs nucleotide bases by the known value for the average spacing between base pairs the nucleotide bases;
 - determining a first resonant frequency of the genomic material to influence the medium-sensitive target nucleic acid chain in one a first electromagnetic frequency range by dividing the velocity of the electromagnetic radiation through emission in the surrounding medium by the wavelength length parameter of the genomic material target nucleic acid chain;
 - multiplying or dividing the first resonant frequency by a factor of a power of two to provide a second resonant frequency in another electromagnetic frequency range;

dividing the number of nucleotide bases into a constant, said constant constituting a shortened version of the aforesaid entire mathematical procedure;

programming the frequency-emitting frequency-capable emission device to emit the second either first or second resonant frequency; and

selectively influencing the target genomic material nucleic acid chain with the first or second resonant frequency in the another electromagnetic frequency range when the frequency-emitting frequency-capable emission device emits the said first or second resonant frequency into the medium surrounding the target genomic material nucleic acid chain.

2. (Currently Amended) The method of claim 1, wherein determining the wavelength length parameter of the genomic material target nucleic acid chain comprises measuring using the known spacing value between adjacent base pairs nucleotide bases and multiplying the number of base pairs nucleotide bases in the genomic material target nucleic acid chain by the measured known spacing value between adjacent base pairs nucleotide bases, and using the resulting value as a wavelength parameter.

3. (Canceled)

4. (Currently Amended) The method of claim 1, wherein the medium surrounding the genomic material is in-vivo tissue having target nucleic acid chain has a unique electrical permittivity, wherein determining the velocity for the propagation of the electromagnetic radiation through emission in the medium surrounding the genomic material target nucleic acid chain comprises relating the unique electrical permittivity of in-vivo tissue to the velocity, obtaining the unique electrical permittivity value for the medium under consideration, and then determining said medium-associated velocity, wherein velocity = $1 / \sqrt{(\epsilon_0 \mu_0)}$, where ϵ_0 is electrical permittivity, and μ_0 is magnetic permeability. $1 / \sqrt{(\epsilon \mu)}$, where ϵ is the electrical permittivity of the medium, and μ is the magnetic permeability of the medium.

5. (Currently amended) The method of claim 4, further comprising the step of determining a refractive index of the electromagnetic ~~radiation through emission in~~ the in-vivo tissue by dividing the speed of light in a vacuum by the speed of light in ~~the~~ in-vivo tissue, wherein dividing one therapeutic resonant frequency determined for the ~~genomic material~~ target nucleic acid chain surrounded by air by the refractive index for in-vivo tissue yields one of the therapeutic resonant frequencies for the ~~genomic material~~ target nucleic acid chain surrounded by in-vivo tissue.

6. (Currently amended) The method of claim 1, further comprising the steps of:
multiplying the first or second resonant frequency ~~in another electromagnetic frequency range~~ by a positive integer to determine harmonic frequencies,
dividing the first or second resonant frequency by a positive integer to determine subharmonic frequencies,
programming the ~~frequency emitting~~ frequency-capable emission device to emit the harmonic ~~and~~ and/or subharmonic frequencies, and
selectively influencing the target ~~genomic material~~ nucleic acid chain with the first or second resonant frequency, ~~in another electromagnetic frequency range and~~ and/or the harmonic ~~and~~ and/or subharmonic frequencies when the ~~frequency emitting~~ frequency-capable emitting device emits the first or second resonant frequency ~~and~~ and/or the harmonic ~~and~~ and/or subharmonic frequencies into the medium surrounding the target ~~genomic material~~ nucleic acid chain.

7. (Currently amended) The method of claim 1, wherein selectively influencing the target ~~genomic material~~ nucleic acid chain comprises debilitating the target ~~genomic material~~ nucleic acid chain.

8. (Currently amended) The method of claim 1, wherein selectively influencing the target ~~genomic material~~ nucleic acid chain comprises stimulating the target ~~genomic material~~ nucleic acid chain.

9. (Currently Amended) The method of claim 1, wherein selectively influencing the target genomic material nucleic acid chain with the first or second resonant frequency comprises selectively influencing genomic material nucleic acid chains present in humans.

10. (Currently Amended) The method of claim 1, wherein selectively influencing the target genomic material nucleic acid chain with the first or second resonant frequency comprises selectively influencing genomic material nucleic acid chains present in animals.

11. (Currently Amended) The method of claim 1, wherein selectively influencing the target genomic material nucleic acid chain with the first or second resonant frequency comprises selectively influencing genomic material nucleic acid chains present in agricultural settings.

12. (Currently Amended) The method of claim 1, wherein selectively influencing the target genomic material nucleic acid chain with the first or second resonant frequency comprises selectively influencing genomic material nucleic acid chains present in water systems.

13. (Currently Amended) The method of claim 1, wherein selectively influencing the target genomic material nucleic acid chain with the first or second resonant frequency comprises selectively influencing genomic material nucleic acid chains present in food processing systems.

14. (Currently amended) The method of claim 1, wherein the medium surrounding the genomic material target nucleic acid chain is in-vivo tissue, further comprising the steps of:

 determining for the genomic material target nucleic acid chain in a medium of air the first resonant frequency in one electromagnetic frequency range and the second resonant frequency in another electromagnetic frequency range, and

multiplying dividing each of the first resonant frequency in one electromagnetic frequency range and the second resonant frequency in another electromagnetic frequency range determined for the genomic material target nucleic acid chain in a medium of air by the square

~~root of two refractive index for in-vivo tissue to yield corresponding resonant frequencies for the genomic material~~ target nucleic acid chain surrounded by in-vivo tissue.

15. (Cancelled)

16. (Cancelled)

17. (Cancelled)

18. (Cancelled)

19. (Cancelled)

20. (Cancelled)

21. (Cancelled)

22. (Cancelled)

23. (Cancelled)

24. (Cancelled)

25. (Cancelled)

26. (Cancelled)

27. (Cancelled)

28. (Cancelled)

29. (Cancelled)